IT IS CLAIMED:

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- 1. A method of detecting each or any of a plurality of known, selected nucleotide target sequences, comprising:
- (a) contacting the target sequences with a set of electrophoretic tag (e-tag) probes, the set comprising j members, and each of said e-tag probes having the form:
 - (D, M_j) N T_j , where
 - (i) D is a detection group comprising a detectable label;
 - (ii) T_j is an oligonucleotide target-binding moiety having a sequence of nucleotides U_i connected by intersubunit linkages $B_{i,\,i+1}$, where i includes all integers from 1 to n, and n is sufficient to allow the moiety to hybridize specifically with a target nucleotide sequence;
 - (iii) N is a nucleotide joined to U₁ in T_i through a nuclease-cleavable bond;
 - (iv) M_j is a mobility modifier having a charge/mass ratio that imparts a unique and known electrophoretic mobility to a corresponding e-tag reporter of the form (D, M_j) N, within a selected range of electrophoretic mobilities with respect to other e-tag reporters of the same form in the probe set, where the e-tag reporter (D, M_j) N does not itself contain nuclease-cleavable bonds; and
 - (v) (D, M_j) includes both D- M_j and M_j D-; said contacting being carried out under conditions that allow hybridization of the target-binding moiety to complementary target sequences,
- (b) treating the hybridized target sequences with a nuclease under conditions effective to cleave target-hybridized probes at their N U_1 linkages, thereby producing a mixture of one or more corresponding e-tag reporters of the form (D, M_j) N, and uncleaved and/or partially cleaved probes,
- (c) exposing the mixture to a capture agent effective to bind to uncleaved and/or partially cleaved probes, but not to e-tag reporters, thereby to (i) impart a mobility to the probes bound to capture agent that prevents the probes from electrophoretically migrating within said range of electrophoretic mobilities or (ii) immobilize the probes on a solid support,
- (d) fractionating e-tag reporters having the form (D, M_j) N by electrophoresis, to effect separation of the e-tag reporters, and
- (e) identifying the electrophoretic mobilities of one or more electrophoretic bands, each band uniquely corresponding to an e-tag reporter that is uniquely assigned to a known target sequence.
- 2. The method of claim 1, wherein each probe has the form $D M_j N T_j$ and the corresponding e-tag reporter has the form $D M_j N$
- 3. The method of claim 1, wherein each probe has the form M_j D N T_j and the corresponding e-tag reporter has the form M_j D N.
 - 4. The method of claim 1, for use in detecting a single nucleotide polymorphism in a target sequence, wherein the oligonucleotide sequence T_j is selected to allow 5'-probe hybridization to the target sequence only if the target sequence contains a designated base at the site of the polymorphism.

- 5. The method of claim 1, wherein at least one nucleotide U_i in the target-binding moiety contains a capture ligand capable of binding specifically to said capture agent, where $i \ge 1$.
- 5 6. The method claim 5, wherein the capture ligand is biotin, and the capture agent is avidin or streptavidin.
 - 7. The method of claim 5, wherein the capture ligand is an antigen and the capture agent is an antibody or antibody fragment that binds specifically to the antigen.
 - 8. The method of claim 1, wherein the capture agent is a polycation and the oligonucleotide has a negatively charged backbone.

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- 9. The method of claim 1, wherein the N U₁ linkage is a phosphodiester bond, and the target-binding moieties contain a nuclease-resistant bond B_{i, i+1}, where i includes at least 1, and the nuclease-resistant bond(s) is one or more linkages selected from the group consisting of thiophosphate, phosphinate, phosphoramidate, amide, and boronate linkages.
- 10. The method of claim 9, wherein at least one nucleotide U_i, i > 1 in said oligonucleotide contains a capture ligand capable of binding specifically to said capture agent.